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What Is Claimed Is:

1. A therapeutic agent for combating Alzheimer's disease, wherein said agent can replace or supplement $\alpha_2 M$ function, or suppress expression of *A2M-2*.

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- 2. An anti-LRP-A β molecule comprising, an A β binding domain, and an LRP binding domain, or a pharmaceutically acceptable salt thereof.
- 3. The anti-LRP-A β molecule of claim 2, wherein said molecule is a peptide, or a pharmaceutically acceptable salt thereof.
 - 4. An anti-LRP-A β peptide comprising:
- (a) an A β binding domain comprising 10-50 contiguous residues of SEQ ID NO:6; and
- (b) an LRP binding domain comprising 10-50 contiguous residues of SEQ ID NO:8, wherein said 10-50 contiguous residues of SEQ ID NO:8 encompass residues 1366-1392, or a pharmaceutically acceptable salt thereof.
 - 5. An anti-LRP-Aβ peptide comprising:
- (a) an A β binding domain having an amino acid sequence selected from the group consisting of SEQ ID NO:6, SEQ ID NO:12, SEQ ID NO:16, SEQ ID NO:18, SEQ ID NO:20, SEQ ID NO:22, SEQ ID NO:24, and SEQ ID NO:26; and
- (b) an LRP binding domain having the amino acid sequence of SEQ ID NO:10, or a pharmaceutically acceptable salt thereof.
 - 6. An anti-LRP-A β peptide comprising:

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(a) an A β binding domain having an amino acid sequence selected from the group consisting of SEQ ID NO:12, SEQ ID NO:16, SEQ

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ID NO:18, SEQ ID NO:20, SEQ ID NO:22, SEQ ID NO:24, and SEQ ID NO:26; and

- (b) an LRP binding domain comprising 10-50 contiguous residues of SEQ ID NO:8, or a pharmaceutically acceptable salt thereof.
- 7. The anti-LRP-A β peptide of claims 4, 5 or 6, wherein said A β binding domain is connected to said LRP binding domain by a peptide bond.
- 8. The anti-LRP-A β peptide of claims 4, 5 or 6, wherein said A β binding domain is connected to said LRP binding domain by a linker.
- 9. The anti-LRP-Aβ peptide of claim 8, wherein said linker is selected from the group consisting of a peptide, or polyethylene glycol.
- 10. The anti-LRP-A β peptide of claims 9, wherein said peptide comprises 1-20 glycine residues.
- 11. A nucleic acid comprising a polynucleotide encoding the anti-LRP-A β peptide of claims 4, 5, 6, 7, 8, 9 or 10.
- 12. An anti-LRP-A β peptide comprising a polypeptide having the sequence of SEQ ID NO:14, or a pharmaceutically acceptable salt thereof.
 - 13. An anti-LRP-Aβ peptide comprising:
- (a) an A β binding domain having an amino acid sequence selected from the group consisting of SEQ ID NO:6, SEQ ID NO:12, SEQ ID NO:16, SEQ ID NO:18, SEQ ID NO:20, SEQ ID NO:22, SEQ ID NO:24, and SEQ ID NO:26;
- (b) an LRP binding domain having the amino acid sequence of SEQ ID NO:10; and

- (c) a linker connecting said $A\beta$ binding domain to said LRP binding domain.
- 14. A nucleic acid molecule comprising a nucleotide encoding the anti-LRP-A β peptide of claims 12 or 13.
- 15. A nucleic acid molecule encoding an anti-LRP-A β peptide comprising:
- (a) a region encoding an A β binding domain, comprising 30-150 contiguous nucleotides of SEQ ID NO:5; and
- (b) a region encoding an LRP binding domain comprising 30-150 contiguous nucleotides of SEQ ID NO:7.
- 16. A nucleic acid molecule encoding an anti-LRP-A β peptide comprising:
- (a) a region encoding an A β binding domain having a nucleotide sequence selected from the group consisting of SEQ ID NO:5, SEQ ID NO:11, SEQ ID NO:15, SEQ ID NO:17, SEQ ID NO:19, SEQ ID NO:21, SEQ ID NO:23, and SEQ ID NO:25; and
- (b) a region encoding an LRP binding domain having the nucleotide sequence of SEQ ID NO:9.
- 17. A nucleic acid molecule encoding an anti-LRP-A β peptide comprising:
- (a) a region encoding an A β binding domain having a nucleotide sequence selected from the group consisting of SEQ ID NO:11, SEQ ID NO:15, SEQ ID NO:17, SEQ ID NO:19, SEQ ID NO:21, SEQ ID NO:23, and SEQ ID NO:25; and
- (b) a region encoding an LRP binding domain comprising 30-150 contiguous nucleotides of SEQ ID NO:7.

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- 18. The nucleic acid molecule of claims 15, 16, or 17, wherein said region encoding said $A\beta$ binding domain is connected to said region encoding said LRP binding domain by a phosphodiester bond.
- 19. The nucleic acid molecule of claims 15, 16 or 17, wherein said region encoding said $A\beta$ binding domain is connected to said region encoding said LRP binding domain by a nucleotide encoding 1-20 glycine residues.
- 20. A nucleic acid molecule comprising, a polynucleotide having at least 95% homology to the nucleic acid molecule of claims 15, 16, 17, 18 or 19.
- 21. A nucleic acid molecule comprising, a first polynucleotide that hybridizes to a second polynucleotide, wherein said second polynucleotide is complementary to the nucleic acid molecule of claims 15, 16, 17, 18 or 19.
- 22. The nucleic acid molecule of claim 21, wherein said first polynucleotide hybridizes to said second polynucleotide under conditions comprising:
- (a) incubating overnight at 42 °C in a solution consisting of 50% formamide, 5x SSC, 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and a 20 μ g/ml denatured, sheared salmon sperm DNA; and
 - (b) washing at 65°C in a solution consisting of 0.1x SSC.
- 23. A nucleic acid molecule comprising a polynucleotide having the nucleotide sequence of SEQ ID NO:13.
- 24. A nucleic acid molecule comprising a polynucleotide having at least 95% identity to the nucleotide sequence of SEQ ID NO:13.

- 25. A nucleic acid molecule comprising a first polynucleotide that hybridizes to a second polynucleotide, wherein said second polynucleotide is complementary to the nucleotide sequence of SEQ ID NO:13.
- 26. The nucleic acid molecule of claim 25, wherein said first polynucleotide hybridizes to said second polynucleotide under conditions comprising:
- (a) incubating overnight at 42 °C in a solution consisting of 50% formamide, 5x SSC, 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and a 20 μ g/ml denatured, sheared salmon sperm DNA; and
 - (b) washing at 65°C in a solution consisting of 0.1x SSC.
- $27. \hspace{0.5cm} \text{A pharmaceutical composition comprising an anti-LRP-A} \\ \text{molecule, and one or more pharmaceutically acceptable carriers.}$
- 28. A pharmaceutical composition comprising the anti-LRP-A β peptide of claims 4, 5, 6, 7, 8, 9, 10 or 13, or a pharmaceutically acceptable salt thereof, and one or more pharmaceutically acceptable carriers.
- 29. A pharmaceutical composition comprising an anti-LRP-A β peptide having an amino acid sequence selected from the group consisting of SEQ ID NO:4 or SEQ ID NO:14, or a pharmaceutically acceptable salt thereof, and one or more pharmaceutically acceptable carriers.
- 30. A method of combating Alzheimer's Disease in a subject comprising administering an anti-LRP-A β molecule.
- 31. The method of claim 30, wherein said anti-LRP-A β molecule is a peptide.

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- 32. A method of combating Alzheimer's Disease in a subject comprising administering the anti-LRP-A β peptide of claims 4, 5, 6, 7, 8, 9, 10 or 13, or a pharmaceutically acceptable salt thereof.
- 33. A method of combating Alzheimer's Disease in a subject comprising administering an anti-LRP-A β peptide having an amino acid sequence selected from the group consisting of SEQ ID NO:4 and SEQ ID NO:14, or a pharmaceutically acceptable salt thereof.
- 34. An *A2M-2* antisense oligonucleotide comprising a nucleotide designed to target *A2M-2* RNA.
- 35. The *A2M-2* antisense oligonucleotide of claim 34, wherein said RNA is hnRNA.
- 36. The A2M-2 antisense oligonucleotide of claim 34, wherein said RNA is mRNA.
- 37. An A2M-2 antisense oligonucleotide comprising a nucleotide having the sequence of SEQ ID NO:27.
- 38. An A2M-2 antisense oligonucleotide comprising a nucleotide having the sequence of the last 15-30 contiguous nucleotides of SEQ ID NO:27.
- 39. An *A2M-2* antisense oligonucleotide comprising nucleotides 36 -50 of SEQ ID NO:27.
 - 40. An *A2M-2* antisense oligonucleotide comprising nucleotides 20 -50 of SEQ ID NO:27.

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- 41. A pharmaceutical composition comprising the *A2M-2* antisense oligonucleotide of claims 34, 35, 36, 37, 38, 39 or 40, and one or more pharmaceutically acceptable carriers.
- 42. A method of combating Alzheimer's Disease in a subject comprising administering the *A2M-2* antisense oligonucleotide of claims 34, 35, 36, 37, 38, 39 or 40.
- 43. A vector for gene therapy of Alzheimer's Disease, comprising a viral vector, wherein said viral vector carries a transgene selected from the group consisting of a gene encoding $\alpha_2 M$, and a gene encoding an anti-LRP-A β peptide.
- 44. The viral vector of claim 43, wherein said transgene is a gene encoding $\alpha_2 M$.
- 45. The viral vector of claim 44, wherein said transgene has the nucleotide sequence of nucleotides 44-4465 of SEQ ID NO:1.
- 46. The viral vector of claim 43, wherein said transgene is a gene encoding an anti-LRP-A β peptide.
- 47. The viral vector of claim 43, where in said transgene encodes the anti-LRP-AB peptide of claims 4, 5, 6, 7, 8, 9, 10, 12 or 13.
- 48. The viral vector of claims 43, 44, 45, 46 or 47, wherein said viral vector is an adeno-associated virus.
- 49. A pharmaceutical composition comprising the viral vector of claims 43, 44, 45, 46, 47 or 48, and one or more pharmaceutically acceptable carriers.

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- 50. A method of combating Alzheimer's Disease in a subject by administering the viral vector of claims 43, 44, 45, 46, 47 or 48.
- 51. A method of screening for a therapeutic agent for Alzheimer's Disease, wherein said therapeutic agent is the agent of claim 1.
- 52. A method of screening for a therapeutic agent for Alzheimer's Disease comprising the steps of:
- (a) incubating cells in the presence of a test agent, wherein said cells are heterozygous or homozygous for the A2M-2 allele, and wherein said cells express A2M-2; and
- (b) determining whether the ratio of normal to aberrant *A2M* mRNA has increased relative to the ratio of normal to aberrant *A2M* mRNA found in cells untreated with test agent.
 - 53. The method of claim 52, wherein said cells are glioma cells.
 - 54. The method of claim 52, wherein said cells are hepatoma cells.
- 55. The method of claim 52, wherein said cells are heterozygous for the A2M-2 allele.
- 56. The method of claim 52, wherein said cells are homozygous for the *A2M-2* allele.
- 57. The method of claim 52 wherein said step (b) comprises S1 nuclease analysis using a probe complementary to SEQ ID NO:1, wherein said probe encompasses nucleotides 2057-2284 of SEQ ID NO:1.
 - 58. The method of claim 57, wherein said probe is 300 bp long.

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- 59. The method of claim 52, wherein said step (b) comprises S1 nuclease analysis using a probe complementary to nucleotides 2024-2323 of SEQ ID NO:1.
- 60. The method of claim 52, wherein said step (b) comprises RT PCR analysis.
- 61. The method of claim 60, wherein said step (b) comprises RT PCR analysis using primers designed to amplify a region of *A2M* encompassing exons 17-18.
- 62. The method of claim 61, wherein said region of *A2M* encompassing exons 17-18 is 300 bp long.
- 63. The method of claim 61, wherein said primers are designed to amplify nucleotides 2052-2289 of SEQ ID NO:1.
- 64. The method of claim 61, wherein said primers consist of a first primer having a nucleotide sequence complementary to nucleotides 2024-2038 of SEQ ID NO:1, and a second primer having the nucleotide sequence of nucleotides 2309-2323 of SEQ ID NO:1.
- 65. A method of screening for a therapeutic agent for Alzheimer's Disease comprising the steps of:
 - (a) incubating $\alpha_2 M$ with a test agent; and
- (b) determining whether said $\alpha_2 M$ of step (b) has undergone a conformational change; wherein said steps are performed in sequential order.
- 66. The method of claim 65, wherein said step (b) comprises performing an $\alpha_2 M$ electrophoretic mobility assay.

A method of screening for a therapeutic agent for Alzheimer's

incubating said well with the substrate for said enzyme;

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Disease comprising the steps of:

(c)

wherein said steps are performed in sequential order.

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incubating α₂M with a test agent; and (a) (b) determining whether said $\alpha_2 M$ of step (b) can bind to LRP; wherein said steps are performed in sequential order. 68. The method of claims 65, 66 or 67, wherein said $\alpha_2 M$ is tetrameric. 69. The method of claim 67, wherein said step (b) comprises performing an ELISA. 70. The method of claim 69, wherein said ELISA comprises the steps of: incubating LRP in a well coated with anti-LRP IgG; (a) incubating said well with said $\alpha_2 M$; (b) (c) incubating said well with anti-α₂M IgG conjugated to an enzyme; and (d) incubating said well with a substrate for said enzyme; wherein said steps are performed in sequential order. 71. The method of claim 69, wherein said ELISA comprises the steps of: (a) incubating a well coated with LRP with said α_2 M; incubating said well with anti-α₂M IgG conjugated to an (b) enzyme; and

- 72. The method of claim 69, wherein said ELISA comprises the steps of:
- (a) incubating said $\alpha_2 M$ in a well coated with an anti- $\alpha_2 M$ IgG specific for activated α_2 M;
 - incubating said well with said $\alpha_2 M$; (b)
- (c) incubating said well with anti-α₂M IgG conjugated to an enzyme; and
- (d) incubating said well with a substrate for said enzyme; wherein said steps are performed in sequential order.
- 73. The method of claim 67, wherein said step (b) comprises immunoblotting.
- 74. The method of claim 73, wherein anti-LRP IgG and anti-α₂M IgG are used to perform said immunoblotting.
- 75. The method of claim 67, wherein said step (b) comprises determining the ability of said α₂M to undergo LRP mediated endocytosis.
- 76. The method of claim 67, wherein said step (b) comprises determining the ability of said α₂M to undergo LRP mediated degradation.